

E. Kommerer

**RAW SEQUENCE LISTING
PATENT APPLICATION US/08/954,954A****1646****Paper No. 8**

DATE: 03/11/1999

TIME: 11:17:34

INPUT SET: S30993.raw

This Raw Listing contains the General
Information Section and up to the first 5 pages.

1 **SEQUENCE LISTING**
2

3 (1) General Information

4
5 (i) APPLICANT: Summers, Neena
6 McWherter, Charles
7 Feng, Yiqing8
9 (ii) TITLE OF THE INVENTION: Novel Erythropoietin Receptor
10 Agonists11
12 (iii) NUMBER OF SEQUENCES: 13513
14 (iv) CORRESPONDENCE ADDRESS:15 (A) ADDRESSEE: G. D. Searle & Co.
16 (B) STREET: P.O. Box 5110
17 (C) CITY: Chicago
18 (D) STATE: IL
19 (E) COUNTRY: U. S. A.
20 (F) ZIP: 6068021
22 (v) COMPUTER READABLE FORM:23 (A) MEDIUM TYPE: Diskette
24 (B) COMPUTER: IBM Compatible
25 (C) OPERATING SYSTEM: DOS
26 (D) SOFTWARE: FastSEQ for Windows Version 2.027
28 (vi) CURRENT APPLICATION DATA:29 (A) APPLICATION NUMBER: 08/954,954
30 (B) FILING DATE: 21-OCT-1997
31 (C) CLASSIFICATION:32
33 (vii) PRIOR APPLICATION DATA:34 (A) APPLICATION NUMBER: 60/034,044
35 (B) FILING DATE: 25-OCT-199636
37
38 (viii) ATTORNEY/AGENT INFORMATION:39 (A) NAME: Bennett, Dennis A
40 (B) REGISTRATION NUMBER: 34,547
41 (C) REFERENCE/DOCKET NUMBER: 2991/142
43 (ix) TELECOMMUNICATION INFORMATION:44 (A) TELEPHONE: 314-737-6986
45 (B) TELEFAX: 314-737-6972**ENTERED**

**RAW SEQUENCE LISTING
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47 (C) TELEX:

48

49

50 (2) INFORMATION FOR SEQ ID NO:1:

51

52 (i) SEQUENCE CHARACTERISTICS:

53 (A) LENGTH: 170 amino acids

54 (B) TYPE: amino acid

55 (C) STRANDEDNESS: single

56 (D) TOPOLOGY: linear

57

58 (ii) MOLECULE TYPE: None

59

60 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

61

62 Asn Ile Thr Thr Gly Cys Ala Glu His Cys Ser Leu Asn Glu Asn Ile
63 1 5 10 15
64 Thr Val Pro Asp Thr Lys Val Asn Phe Tyr Ala Trp Lys Arg Met Glu
65 20 25 30
66 Val Gly Gln Gln Ala Val Glu Val Trp Gln Gly Leu Ala Leu Leu Ser
67 35 40 45
68 Glu Ala Val Leu Arg Gly Gln Ala Leu Leu Val Asn Ser Ser Gln Pro
69 50 55 60
70 Trp Glu Pro Leu Gln Leu His Val Asp Lys Ala Val Ser Gly Leu Arg
71 65 70 75 80
72 Ser Leu Thr Thr Leu Leu Arg Ala Leu Gly Ala Gln Lys Glu Ala Ile
73 85 90 95
74 Ser Pro Pro Asp Ala Ala Ser Ala Ala Pro Leu Arg Thr Ile Thr Ala
75 100 105 110
76 Asp Thr Phe Arg Lys Leu Phe Arg Val Tyr Ser Asn Phe Leu Arg Gly
77 115 120 125
78 Lys Leu Lys Leu Tyr Thr Gly Glu Ala Cys Arg Thr Gly Asp Arg Gly
79 130 135 140
80 Gly Gly Ser Ala Pro Pro Arg Leu Ile Cys Asp Ser Arg Val Leu Glu
81 145 150 155 160
82 Arg Tyr Leu Leu Glu Ala Lys Glu Ala Glu
83 165 170

84

85 (2) INFORMATION FOR SEQ ID NO:2:

86

87 (i) SEQUENCE CHARACTERISTICS:

88 (A) LENGTH: 170 amino acids

89 (B) TYPE: amino acid

90 (C) STRANDEDNESS: single

91 (D) TOPOLOGY: linear

92

93 (ii) MOLECULE TYPE: None

94

95 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

96

97 Ile Thr Thr Gly Cys Ala Glu His Cys Ser Leu Asn Glu Asn Ile Thr
98 1 5 10 15
99 Val Pro Asp Thr Lys Val Asn Phe Tyr Ala Trp Lys Arg Met Glu Val

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100      20          25          30
101 Gly Gln Gln Ala Val Glu Val Trp Gln Gly Leu Ala Leu Leu Ser Glu
102      35          40          45
103 Ala Val Leu Arg Gly Gln Ala Leu Leu Val Asn Ser Ser Gln Pro Trp
104      50          55          60
105 Glu Pro Leu Gln Leu His Val Asp Lys Ala Val Ser Gly Leu Arg Ser
106      65          70          75          80
107 Leu Thr Thr Leu Leu Arg Ala Leu Gly Ala Gln Lys Glu Ala Ile Ser
108      85          90          95
109 Pro Pro Asp Ala Ala Ser Ala Ala Pro Leu Arg Thr Ile Thr Ala Asp
110      100         105         110
111 Thr Phe Arg Lys Leu Phe Arg Val Tyr Ser Asn Phe Leu Arg Gly Lys
112      115         120         125
113 Leu Lys Leu Tyr Thr Gly Glu Ala Cys Arg Thr Gly Asp Arg Gly Gly
114      130         135         140
115 Gly Ser Ala Pro Pro Arg Leu Ile Cys Asp Ser Arg Val Leu Glu Arg
116      145         150         155         160
117 Tyr Leu Leu Glu Ala Lys Glu Ala Glu Asn
118      165         170
119
120      (2) INFORMATION FOR SEQ ID NO:3:
121
122      (i) SEQUENCE CHARACTERISTICS:
123          (A) LENGTH: 170 amino acids
124          (B) TYPE: amino acid
125          (C) STRANDEDNESS: single
126          (D) TOPOLOGY: linear
127
128      (ii) MOLECULE TYPE: None
129
130      (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:
131
132 Thr Thr Gly Cys Ala Glu His Cys Ser Leu Asn Glu Asn Ile Thr Val
133      1           5           10          15
134 Pro Asp Thr Lys Val Asn Phe Tyr Ala Trp Lys Arg Met Glu Val Gly
135      20          25          30
136 Gln Gln Ala Val Glu Val Trp Gln Gly Leu Ala Leu Leu Ser Glu Ala
137      35          40          45
138 Val Leu Arg Gly Gln Ala Leu Leu Val Asn Ser Ser Gln Pro Trp Glu
139      50          55          60
140 Pro Leu Gln Leu His Val Asp Lys Ala Val Ser Gly Leu Arg Ser Leu
141      65          70          75          80
142 Thr Thr Leu Leu Arg Ala Leu Gly Ala Gln Lys Glu Ala Ile Ser Pro
143      85          90          95
144 Pro Asp Ala Ala Ser Ala Ala Pro Leu Arg Thr Ile Thr Ala Asp Thr
145      100         105         110
146 Phe Arg Lys Leu Phe Arg Val Tyr Ser Asn Phe Leu Arg Gly Lys Leu
147      115         120         125
148 Lys Leu Tyr Thr Gly Glu Ala Cys Arg Thr Gly Asp Arg Gly Gly Gly
149      130         135         140
150 Ser Ala Pro Pro Arg Leu Ile Cys Asp Ser Arg Val Leu Glu Arg Tyr
151      145         150         155         160
152 Leu Leu Glu Ala Lys Glu Ala Glu Asn Ile

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**RAW SEQUENCE LISTING
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153 165 170

154
155 (2) INFORMATION FOR SEQ ID NO:4:

156

157 (i) SEQUENCE CHARACTERISTICS:

158 (A) LENGTH: 170 amino acids

159 (B) TYPE: amino acid

160 (C) STRANDEDNESS: single

161 (D) TOPOLOGY: linear

162

163 (ii) MOLECULE TYPE: None

164

165 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

166

167 Thr	Gly	Cys	Ala	Glu	His	Cys	Ser	Leu	Asn	Glu	Asn	Ile	Thr	Val	Pro
168 1	5							10					15		
169 Asp	Thr	Lys	Val	Asn	Phe	Tyr	Ala	Trp	Lys	Arg	Met	Glu	Val	Gly	Gln
170 20		20					25					30			
171 Gln	Ala	Val	Glu	Val	Trp	Gln	Gly	Leu	Ala	Leu	Leu	Ser	Glu	Ala	Val
172 35		35					40					45			
173 Leu	Arg	Gly	Gln	Ala	Leu	Leu	Val	Asn	Ser	Ser	Gln	Pro	Trp	Glu	Pro
174 50		50					55				60				
175 Leu	Gln	Leu	His	Val	Asp	Lys	Ala	Val	Ser	Gly	Leu	Arg	Ser	Leu	Thr
176 65		65				70		75			80				
177 Thr	Leu	Leu	Arg	Ala	Leu	Gly	Ala	Gln	Lys	Glu	Ala	Ile	Ser	Pro	Pro
178 85		85						90			95				
179 Asp	Ala	Ala	Ser	Ala	Ala	Pro	Leu	Arg	Thr	Ile	Thr	Ala	Asp	Thr	Phe
180 100		100						105			110				
181 Arg	Lys	Leu	Phe	Arg	Val	Tyr	Ser	Asn	Phe	Leu	Arg	Gly	Lys	Leu	Lys
182 115		115					120			125					
183 Leu	Tyr	Thr	Gly	Glu	Ala	Cys	Arg	Thr	Gly	Asp	Arg	Gly	Gly	Gly	Ser
184 130		130					135			140					
185 Ala	Pro	Pro	Arg	Leu	Ile	Cys	Asp	Ser	Arg	Val	Leu	Glu	Arg	Tyr	Leu
186 145		145				150			155			160			
187 Leu	Glu	Ala	Lys	Glu	Ala	Glu	Asn	Ile	Thr						
188 165		165						170							

189

190 (2) INFORMATION FOR SEQ ID NO:5:

191

192 (i) SEQUENCE CHARACTERISTICS:

193 (A) LENGTH: 170 amino acids

194 (B) TYPE: amino acid

195 (C) STRANDEDNESS: single

196 (D) TOPOLOGY: linear

197

198 (ii) MOLECULE TYPE: None

199

200 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

201

202 Gly	Cys	Ala	Glu	His	Cys	Ser	Leu	Asn	Glu	Asn	Ile	Thr	Val	Pro	Asp
203 1	5							10				15			
204 Thr	Lys	Val	Asn	Phe	Tyr	Ala	Trp	Lys	Arg	Met	Glu	Val	Gly	Gln	Gln
205 20		20						25			30				

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206 Ala Val Glu Val Trp Gln Gly Leu Ala Leu Leu Ser Glu Ala Val Leu
207      35          40          45
208 Arg Gly Gln Ala Leu Leu Val Asn Ser Ser Gln Pro Trp Glu Pro Leu
209      50          55          60
210 Gln Leu His Val Asp Lys Ala Val Ser Gly Leu Arg Ser Leu Thr Thr
211      65          70          75          80
212 Leu Leu Arg Ala Leu Gly Ala Gln Lys Glu Ala Ile Ser Pro Pro Asp
213      85          90          95
214 Ala Ala Ser Ala Ala Pro Leu Arg Thr Ile Thr Ala Asp Thr Phe Arg
215      100         105         110
216 Lys Leu Phe Arg Val Tyr Ser Asn Phe Leu Arg Gly Lys Leu Lys Leu
217      115         120         125
218 Tyr Thr Gly Glu Ala Cys Arg Thr Gly Asp Arg Gly Gly Ser Ala
219      130         135         140
220 Pro Pro Arg Leu Ile Cys Asp Ser Arg Val Leu Glu Arg Tyr Leu Leu
221      145         150         155         160
222 Glu Ala Lys Glu Ala Glu Asn Ile Thr Thr
223      165         170
224
225 (2) INFORMATION FOR SEQ ID NO:6:
226
227 (i) SEQUENCE CHARACTERISTICS:
228   (A) LENGTH: 170 amino acids
229   (B) TYPE: amino acid
230   (C) STRANDEDNESS: single
231   (D) TOPOLOGY: linear
232
233 (ii) MOLECULE TYPE: None
234
235 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:
236
237 Cys Ala Glu His Cys Ser Leu Asn Glu Asn Ile Thr Val Pro Asp Thr
238      1          5          10          15
239 Lys Val Asn Phe Tyr Ala Trp Lys Arg Met Glu Val Gly Gln Gln Ala
240      20         25         30
241 Val Glu Val Trp Gln Gly Leu Ala Leu Leu Ser Glu Ala Val Leu Arg
242      35         40         45
243 Gly Gln Ala Leu Leu Val Asn Ser Ser Gln Pro Trp Glu Pro Leu Gln
244      50         55         60
245 Leu His Val Asp Lys Ala Val Ser Gly Leu Arg Ser Leu Thr Thr Leu
246      65         70         75          80
247 Leu Arg Ala Leu Gly Ala Gln Lys Glu Ala Ile Ser Pro Pro Asp Ala
248      85         90         95
249 Ala Ser Ala Ala Pro Leu Arg Thr Ile Thr Ala Asp Thr Phe Arg Lys
250      100        105        110
251 Leu Phe Arg Val Tyr Ser Asn Phe Leu Arg Gly Lys Leu Lys Leu Tyr
252      115        120        125
253 Thr Gly Glu Ala Cys Arg Thr Gly Asp Arg Gly Gly Ser Ala Pro
254      130        135        140
255 Pro Arg Leu Ile Cys Asp Ser Arg Val Leu Glu Arg Tyr Leu Leu Glu
256      145        150        155        160
257 Ala Lys Glu Ala Glu Asn Ile Thr Thr Gly
258      165        170

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**SEQUENCE VERIFICATION REPORT
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Line

Error

Original Text